

Triple knockout of *frdC* *gltA* and *pta* genes enhanced PHA production in *Escherichia coli*

ABSTRACT

Polyhydroxyalkanoate (PHA) is a linear polyester produced through the fermentation of sugar or lipid. Biosynthesis of PHA comprises three enzymes known as acetyl-CoA acetyltransferase (*phaA*), acetoacetyl-CoA reductase (*phaB*) and PHA synthase (*phaC*). *Comamonas* sp. is one of the strains commonly used for PHA production. In order to develop higher PHA production from bacterial respond strategy, PHA biosynthesis operon of *Comamonas* sp. EB172 was introduced into *Escherichia coli* BW25113 through a pGEM-T vector. *E. coli* was chosen due to the complete genome information available and the absence of depolymerisation gene, *phaZ*. In this study, the deletion of several single genes, which are *frdC*, *gltA*, and *pta*, was found to be associated with PHA metabolism activity in *E. coli* BW25113. P1 transduction was performed to construct multiple genes knockout. The engineered strain, *E. coli* BW25113 *frdCgltApta::kan/pGEM⁺-phaCABC*_{Co}, yielded the highest PHA production at 64 wt.% with 1.4 fold higher than that of control strain of *E. coli* BW25113/pGEM⁺-*phaCABC*_{Co}. This strain is potential for industrial application for higher PHA production from *E. coli*.

Keyword: *Escherichia coli*; *FrdC*; *GltA*; P1 transduction; Polyhydroxyalkanoates; *Pta*